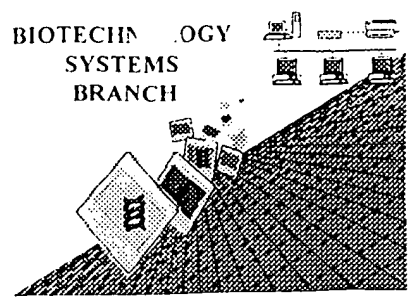


0400



RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/765,614
Source: OIPE
Date Processed by STIC: 2-5-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/765,614

DATE: 02/05/2001
TIME: 13:37:22

Input Set : A:\seqlisting7-10-00.txt
Output Set: N:\CRF3\02052001\I765614.raw

**Does Not Comply
Corrected Diskette Needed**

5 <110> APPLICANT: Nycomed Imaging AS
7 <120> TITLE OF INVENTION: Improvements in or relating to diagnostic/therapeutic
8 agents
10 <130> FILE REFERENCE: REF/Klaveness/054
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/765,614
C--> 13 <141> CURRENT FILING DATE: 2001-01-22
E--> 15 <160> NUMBER OF SEQ ID NOS: 25
17 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

Number of sequences differ:
- 25 listed
- 31 shown (see next page)

09/765, 614

p. 2

<210> 31
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Platelet
binding peptide

Last sequence

<400> 31
Pro Leu Tyr Lys Lys Ile Ile Lys Lys Leu Leu Glu Ser
1 5 10

in file, #31.

VERIFICATION SUMMARY

DATE: 02/05/2001

PATENT APPLICATION: US/09/765,614

TIME: 13:37:23

Input Set : A:\seqlisting7-10-00.txt

Output Set: N:\CRF3\02052001\I765614.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:301 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:15 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (25) Counted (31)